

**Table II**  
**Physical characteristics of 3'-O-(2-methoxyethyl)**  
**containing 2'-5' linked oligonucleotides.**

|              | Expected Mass | Observed Mass | HPLC <sup>2</sup> T <sub>R</sub> Purified | #Ods(260nm)<br>(min.) |
|--------------|---------------|---------------|---|-----------------------|
| <b>17176</b> | 6440.743      | 6440.300      | 23.47                                     | 3006                  |
| <b>17177</b> | 6514.814      | 6513.910      | 23.67                                     | 3330                  |
| <b>17178</b> | 6482.814      | 6480.900      | 23.06                                     | 390                   |
| <b>17179</b> | 6513.798      | 6513.560      | 23.20                                     | 3240                  |
| <b>17180</b> | 6588.879      | 6588.200      | 23.96                                     | 3222                  |
| <b>17181</b> | 6540.879      | 6539.930      | 23.01                                     |                       |
| <b>21415</b> | 6662.976      | 6662.700      | 24.18                                     | 4008                  |
| <b>21416</b> | 6598.969      | 6597.800      | 23.01                                     | 3060                  |
| <b>21945</b> | 1099.924      | 1099.300      | 19.92                                     | 121                   |
| <b>21663</b> | 1487.324      | 1486.800      | 20.16                                     | 71                    |
| <b>20389</b> | 1483.000      | 1482.000      |   | 62                    |
| <b>20390</b> | 4588.000      | 4591.000      |   | 151                   |

<sup>2</sup>Conditions: Waters 600E with detector 991; Waters C4 column (3.9X300mm); Solvent A: 50 mM TEA-Ac, pH 7.0; B: 100% acetonitrile; 1.5 mL/min. flow rate; Gradient: 5% B for first five minutes with linear increase in B to 60% during the next 55 minutes.

## EXAMPLE 51

### **T<sub>m</sub> Studies on modified oligonucleotides**

[0188] Oligonucleotides synthesized in Examples 49 and 50 were evaluated for their relative ability to bind to their complementary nucleic acids by measurement of their melting temperature (T<sub>m</sub>). The melting temperature (T<sub>m</sub>), a characteristic physical property of double helices, denotes the temperature (in degrees centigrade) at which 50% helical (hybridized) versus coil (unhybridized) forms are present. T<sub>m</sub> is measured by using the UV spectrum to determine the formation and breakdown (melting) of the hybridization complex. Base stacking, which occurs during hybridization, is accompanied by a reduction in UV absorption (hypochromicity). Consequently, a reduction in UV absorption indicates a higher T<sub>m</sub>. The higher the T<sub>m</sub>, the greater the strength of the bonds between the strands.

[0189] Selected test oligonucleotides and their complementary nucleic acids were incubated at a standard concentration of 4  $\mu$ M for each oligonucleotide in buffer (100 mM NaCl, 10 mM sodium phosphate, pH 7.0, 0.1 mM EDTA). Samples were heated to 90 °C and the initial absorbance taken using a Guilford Response II Spectrophotometer (Corning). Samples were then slowly cooled to 15 °C and then the change in absorbance at 260 nm was monitored with

heating during the heat denaturation procedure. The temperature was increased by 1 degree °C/absorbance reading and the denaturation profile analyzed by taking the 1<sup>st</sup> derivative of the melting curve. Data was also analyzed using a two-state linear regression analysis to determine the Tm=s. The results of these tests for the some of the oligonucleotides from Examples 49 and 50 are shown in Table III below.

Table III

| Tm Analysis of Oligonucleotides |  |          |                |   |                    |
|---------------------------------|--|----------|----------------|---|--------------------|
| SEQ ID:<br>NO. #                | (ISIS) # Sequence (5'-3')  | Backbone | T <sub>m</sub> | 8 | # Mods<br>Linkages |
| 13                              | (11061) ATG-CAT-TCT-GCC-CCC-AAG-GA   | P=S      | 61.4           | 0 | 0                  |
| 4                               | (17176) ATG-CAT-TCT-GCC-CCC-AAG-GA*  | P=S      | 61.4           | 1 | 0                  |
| 5                               | (17177) ATG-CAT-TCT-GCC-CCC-AAG-G*A*   | P=S      | 61.3           | 2 | 1                  |
| 6                               | (17178) ATG-CAT-TCT-GCC-CCC-AAG <sub>o</sub> -G* <sub>o</sub> A*                               | P=S/P=O  | 61.8           | 2 | 1                  |
| 7                               | (17179) A*TG-CAT-TCT-GCC-CCC-AAG-GA*   | P=S      | 61.1           | 2 | 1                  |
| 8                               | (17180) A*TG-CAT-TCT-GCC-CCC-AAG-G*A*  | P=S      | 61.0           | 3 | 2                  |
| 9                               | (17181) A* <sub>o</sub> TG-CAT-TCT-GCC-AAA-AAG <sub>o</sub> -G* <sub>o</sub> A*                | P=S/P=O  | 61.8           | 3 | 2                  |
| 10                              | (21415) A*T*G-CAT-TCT-GCC-AAA-AAG-G*A*   | P=S      | 61.4           | 4 | 3                  |
| 11                              | (21416) A* <sub>o</sub> T* <sub>o</sub> G-CAT-TCT-GCC-AAA-AAG <sub>o</sub> -G* <sub>o</sub> A* | P=S/P=O  | 61.7           | 4 | 3                  |